/FWO

CRF Errors Edited by the STIC Systems Branch

1 Number: <u>/0/1/06,43</u>	CRF Edit Date: 9/29/0 Edited by:
Realigned nucleic acid/a find text in the mext in	acid numbers/text in cases where the sequence
Corrected the SEQ ID NO. S	Sequence numbers edited were:
Inserted or corrected a nucle NO's edited:	cic number at the end of a nucleic line. SEQ ID
	•
Deleted: invalid beginni	ng/end-of-file text; page numbers
Inserted mandatory heading	s/numeric identifiers, specifically:
Moved responses to same lin	e as heading/numeric identifier, specifically:
	ted anew aid humbering; Inserte numeric identifier after 4th line



IFWO

RAW SEQUENCE LISTINGPATENT APPLICATION: **US/10/706,435A**DATE: 09/29/2004
TIME: 11:12:57

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\09292004\J706435A.raw

```
3 <110> APPLICANT: Walter Reed Army Institute of Research
        Lanar, David E.
        Hillier, Collette J.
 5
        Lyon, Jeffrey A.
 6
 7
         Angov, Evelina
 8
         Kumar, Sanjai
9
         Rogers, William
10
         Barbosa, Arnoldo
12 <120> TITLE OF INVENTION: Expression, Purification, and Uses of a Plasmodium
        falciparum Liver Stage Antigen 1 Polypeptide
15 <130> FILE REFERENCE: 003/285/SAP
17 <140> CURRENT APPLICATION NUMBER: 10/706,435A
19 <141> CURRENT FILING DATE: 2003-11-12
21 <150> PRIOR APPLICATION NUMBER: 60/425,719
23 <151> PRIOR FILING DATE: 2002-11-12
25 <160> NUMBER OF SEQ ID NOS: 28
27 <170> SOFTWARE: Microsoft Word XP
29 <210> SEQ ID NO: 1
30 <211> LENGTH: 17
31 <212> TYPE: PRT
32 <213> ORGANISM: P. falciparum LSA-1
33 <220> FEATURE:
34 <223> OTHER INFORMATION: LSA-1 major 17 amino acid repeat
35 <400> SEQUENCE: 1
37 Glu Gln Gln Ser Asp Leu Glu Gln Glu Arg
                    5
39 Leu Ala Lys Glu Lys Leu Gln
40
42 <210> SEQ ID NO: 2
43 <211> LENGTH: 17
44 <212> TYPE: PRT
45 <213> ORGANISM: P. falciparum LSA-1
46 <220> FEATURE:
47 <223> OTHER INFORMATION: LSA-1 minor 17 amino acid repeat
48 <400> SEQUENCE: 2
50 Glu Gln Gln Arg Asp Leu Glu Gln Glu Arg
52 Leu Ala Lys Glu Lys Leu Gln
53
55 <210> SEQ ID NO: 3
56 <211> LENGTH: 1374
57 <212> TYPE: DNA
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58 <213> ORGANISM: Artificial sequence

RAW SEQUENCE LISTING

DATE: 09/29/2004 TIME: 11:12:57

Input Set : A:\PTO.AMC.txt

PATENT APPLICATION: US/10/706,435A

Output Set: N:\CRF4\09292004\J706435A.raw

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59 <220> FEATURE:
60 <223> OTHER INFORMATION: LSA-NRC(H) Mut
61 <400> SEQUENCE: 3
63 atgggtacca acagcgaaaa agacgaaatt atcaaaagca
64 atetecgete eggeagetee aacageegea acegeateaa
65 cqaqqaaaag catqagaaga aacatqtgct gagccacaac
                                                 120
66 tectaegaga agaetaaaaa caaegaaaac aacaaattet
                                                 160
67 ttgacaagga caaagagctg acgatgagca acgttaaaaa
                                                 200
68 cgtatcccag accaacttta aatccctcct gcgcaacctc
                                                 280
69 ggcgtttccg agaacatctt tctcaaagaa aacaaactga
                                                 320
70 acaaggaagg caaactgatt gaacatatca tcaacgacga
                                                 360
71 cqatgacaaa aaaaaataca ttaaaggcca ggatgaaaat
72 cgccaggaag acctcgaaga aaaagctgct gaacagcagt
                                                 400
73 cggacctgga acaggagcgc ctcgctaaag aaaagctcca
74 ggagcgcctc gctaaagaaa agctccagga gcaacagcgc
                                                 480
75 gacctggaac agcgcaaggc tgacacgaaa aaaaacctgg
                                                 520
76 aacgcaaaaa ggaacacggc gacgttctgg ctgaggacct
                                                 560
77 gtacggccgc ctggaaatcc cagctatcga actcccatcc
                                                 600
78 gaaaacgaac gcggctacta catcccacac cagagcagcc
79 tgccacaaga taatcgcggg aactcccgcg acagtaagga
                                                 680
80 aatcagcatc atcgaaaaaa ccaaccgcga aagcattacc
                                                 720
81 accaacgtgg aaggccgccg cgacatccac aaaggccacc
                                                 760
82 tegaagaaaa gaaagaegge teeateaaac cagaacagaa
83 agaagacaaa agcgctgata tccagaacca caccctggag
84 accgtgaaca ttagcgacgt gaacgacttc cagatcagca
85 agtacgágga cgaaatctcc gctgaatacg atgactccct
                                                 920
86 gatcgacgaa gaagaagacg acgaagatct ggatgaattc
87 aaaccaattg tccaqtacqa taactttcaq qacqaaqaaa 1000
88 atateggeat ttacaaagaa etegaagaee teategagaa 1040
89 aaacgaaaac ctggacgacc tggacgaagg catcgaaaaa 1080
90 tcctccgaag aactgagcga agaaaaaatc aaaaaaggca 1120
91 agaaatacga aaaaaccaag gacaacaact tcaaaccaaa 1160
92 cgacaaatcc ctctacgacg agcacattaa aaaatacaaa 1200
93 aacgacaagc aagtgaacaa ggaaaaggaa aaatttatca 1240
94 aatccctctt ccacatcttc gatggcgata acgaaattct 1280
95 gcaaattgta gacgaacggt tgagcgaaga catcactaaa 1320
96 tacttcatga agettggggg ctccggttct ccacaccacc 1360
97 accaccacca ctga
                                                1374
100 <210> SEQ ID NO: 4
101 <211> LENGTH: 457
102 <212> TYPE: PRT
103 <213> ORGANISM: Artificial sequence
104 <220> FEATURE:
105 <223 > OTHER INFORMATION: LSA-NRC(H) Mut
106 <400> SEQUENCE: 4
108 Met Gly Thr Asn Ser Glu Lys Asp Glu Ile
                      5
110 Ile Lys Ser Asn Leu Arg Ser Gly Ser Ser
111
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/706,435A

DATE: 09/29/2004 TIME: 11:12:57

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\09292004\J706435A.raw

	Asn	Ser	Arg	Asn	Arg 25	Ile	Asn	Glu	Glu	Lys 30
113		~ 3		~		**- 7	.	G	***	
114	His	GIU	Lys	ьys		Val	ьeu	Ser	His	Asn
115	_	_		_	35	_	_	_	~ 3	40
116	Ser	Tyr	Glu	Lys	Thr	гàг	Asn	Asn	Glu	Asn
117					45	_	_			50
118	Asn	Lys	Phe	Phe	_	Lys	Asp	Lys	GIu	Leu
119					55					60
120	Thr	Met	Ser	Asn	Val	Lys	Asn	Val	Ser	Gln
121					65					70
122	Thr	Asn	Phe	Lys		Leu	Leu	Arg	Asn	Leu
123					75					80
124	Gly	Val	Ser	Glu	Asn	Ile	Phe	Leu	Lys	Glu
125					85					90
126	Asn	Lys	Leu	Asn	Lys	Glu	Gly	Lys	Leu	Ile
127					95					100
128	Glu	His	Ile	Ile	Asn	qaA	Asp	Asp	Asp	Lys
129					105					110
130	Lys	Lys	Tyr	Ile	Lys	Gly	Gln	Asp	Glu	Asn
131					115					120
132	Arg	Gln	Glu	Asp	Leu	Glu	Glu	Lys	Ala	Ala
133					125					130
134	Glu	Gln	Gln	Ser	Asp	Leu	Glu	Gln	Glu	Arg
135					135					140
136	Leu	Ala	Lys	Glu	Lys	Leu	Gln	Glu	Arg	Leu
137					145					150
138	Ala	Lys	Glu	Lys	Leu	Gln	Glu	Gln	Gln	Arg
139					155		٠			160
140	Asp	Leu	Glu	Gln	Arg	Lys	Ala	Asp	Thr	Lys
141					165					170
142	Lys	Asn	Leu	Glu	Arg	Lys	Lys	Glu	His	Gly
143					175					180
144	Asp	Val	Leu	Ala	Glu	Asp	Leu	Tyr	Gly	Arg
145					185					190
146	Leu	Glu	Ile	Pro	Ala	Ile	Glu	Leu	Pro	Ser
147					195					200
148	Glu	Asn	Glu	Arg	Gly	Tyr	Tyr	Ile	Pro	His
149					205					210
150	Gln	Ser	Ser	Leu	Pro	Gln	Asp	Asn	Arg	Gly
151					215		_			220
152	Asn	Ser	Arq	Asp	Ser	Lys	Glu	Ile	Ser	Ile
153				_	225	_				230
154	Ile	Glu	Lys	Thr	Asn	Arq	Glu	Ser	Ile	Thr
155			4		235	5				240
156	Thr	Asn	Val	Glu	Gly	Ara	Ara	Asp	Ile	
157					245	ر		Ľ	_	250
158	Lvs	Glv	His	Leu	Glu	Glu	Lvs	Lvs	Asp	
159	-10	1			255			-1-		260
162	Ser	Ile	Lvs	Pro	Glu	Gln	Lvs	Glu	Asp	
										4

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/706,435A

DATE: 09/29/2004 TIME: 11:12:57

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\09292004\J706435A.raw

```
163
                     265
164 Ser Ala Asp Ile Gln Asn His Thr Leu Glu
                     275
166 Thr Val Asn Ile Ser Asp Val Asn Asp Phe
167
                     285
                                          290
    Gln Ile Ser Lys Tyr Glu Asp Glu Ile Ser
168
169
                     295
170 Ala Glu Tyr Asp Asp Ser Leu Ile Asp Glu
171
                     305
172 Glu Glu Asp Asp Glu Asp Leu Asp Glu Phe
173
                     315
174 Lys Pro Ile Val Gln Tyr Asp Asn Phe Gln
175
                     325
                                          330
176 Asp Glu Glu Asn Ile Gly Ile Tyr Lys Glu
177
                     335
178 Leu Glu Asp Leu Ile Glu Lys Asn Glu Asn
179
                     345
180 Leu Asp Asp Leu Asp Glu Gly Ile Glu Lys
181
                     355
182 Ser Ser Glu Glu Leu Ser Glu Glu Lys Ile
183
                     365
184 Lys Lys Gly Lys Lys Tyr Glu Lys Thr Lys
                     375
186 Asp Asn Asn Phe Lys Pro Asn Asp Lys Ser
187
                     385
188 Leu Tyr Asp Glu His Ile Lys Lys Tyr Lys
189
                     395
                                          400
190 Asn Asp Lys Gln Val Asn Lys Glu Lys Glu
191
                     405
192 Lys Phe Ile Lys Ser Leu Phe His Ile Phe
193
                     415
194 Asp Gly Asp Asn Glu Ile Leu Gln Ile Val
195
                     425
196 Asp Glu Arg Leu Ser Glu Asp Ile Thr Lys
197
                     435
                                          440
198 Tyr Phe Met Lys Leu Gly Gly Ser Gly Ser
                     445
                                         450
200 Pro His His His His His
201
203 <210> SEQ ID NO: 5
204 <211> LENGTH: 17
205 <212> TYPE: PRT
206 <213> ORGANISM: Artificial sequence
207 <220> FEATURE:
208 <223> OTHER INFORMATION: LSA-1 Consensus sequence of 17 amino acid repeats
209
          where Xaa at position 1 is either Glu or Gly; Xaa at
210
          position 4 is Ser or Arg; Xaa at position 6 is Asp or Ser;
211
          Xaa at position 9 is Glu or Asp; Xaa at position 11 is Leu
212 <220> FEATURE:
```

DATE: 09/29/2004

TIME: 11:12:57

Input Set : A:\PTO.AMC.txt Output Set: N:\CRF4\09292004\J706435A.raw 213 <223> OTHER INFORMATION: or Arg; Xaa at position 13 is Lys or Asn and Xaa at position 214 15 is Lys or Thr or Arq. 216 <400> SEQUENCE: 5 W--> 217 Xaa Gln Gln Xaa Asp Xaa Glu Gln Xaa Arg 218 220 Xaa Ala Xaa Glu Xaa Leu Gln 221 15 223 <210> SEQ ID NO: 6 224 <211> LENGTH: 24 225 <212> TYPE: PRT 226 <213> ORGANISM: P. falciparum LSA-1 227 <220> FEATURE: 228 <223> OTHER INFORMATION: P. falciparum LSA-1 T1 epitope 229 <400> SEQUENCE: 6 230 Leu Thr Met Ser Asn Val Lys Asn Val Ser 231 5 232 Gln Thr Asn Phe Lys Ser Leu Leu Arg Asn 233 15 234 Leu Gly Val Ser 236 <210> SEQ ID NO: 7 237 <211> LENGTH: 17 238 <212> TYPE: PRT 239 <213> ORGANISM: P. falciparum LSA-1 240 <220> FEATURE: 241 <223> OTHER INFORMATION: P. falciparum LSA-1 LSA-Rep epitope 242 <400> SEQUENCE: 7 243 Glu Gln Gln Ser Asp Leu Glu Gln Glu Arg 245 Leu Ala Lys Glu Lys Leu Gln 246 248 <210> SEQ ID NO: 8 249 <211> LENGTH: 17 250 <212> TYPE: PRT 251 <213> ORGANISM: P. falciparum LSA-1 252 <220> FEATURE: 253 <223> OTHER INFORMATION: P. falciparum LSA-1 J epitope 254 <400> SEQUENCE: 8 255 Glu Arg Leu Ala Lys Glu Lys Leu Gln Glu 10 257 Gln Gln Arg Asp Leu Glu Gln 258 260 <210> SEQ ID NO: 9 261 <211> LENGTH: 20 262 <212> TYPE: PRT 263 <213> ORGANISM: P. falciparum LSA-1 264 <220> FEATURE: 265 <223> OTHER INFORMATION: P. falciparum LSA-1 NR epitope 266 <400> SEQUENCE: 9 269 Thr Lys Lys Asn Leu Glu Arg Lys Lys Glu

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/706,435A

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/706,435A

DATE: 09/29/2004 TIME: 11:12:58

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\09292004\J706435A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:5; Xaa Pos. 1,4,6,9,11,13,15



IFWO

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/706,435A

DATE: 09/27/2004 TIME: 11:39:09

Input Set : A:\PTO.LM.txt

Output Set: N:\CRF4\09272004\J706435A.raw

```
3 <110> APPLICANT: Walter Reed Army Institute of Research
 4
         Lanar, David E.
 5
         Hillier, Collette J.
         Lyon, Jeffrey A.
 7
         Angov, Evelina
         Kumar, Sanjai
 8
         Rogers, William
 9
10
         Barbosa, Arnoldo
12 <120> TITLE OF INVENTION: Expression, Purification, and Uses of a Plasmodium
         falciparum Liver Stage Antigen 1 Polypeptide
13
15 <130> FILE REFERENCE: 003/285/SAP
17 <140> CURRENT APPLICATION NUMBER: 10/706,435A
19 <141> CURRENT FILING DATE: 2003-11-12
21 <150> PRIOR APPLICATION NUMBER: 60/425,719
23 <151> PRIOR FILING DATE: 2002-11-12
25 <160> NUMBER OF SEQ ID NOS: 28
27 <170> SOFTWARE: Microsoft Word XP
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ERRORED SEQUENCES

203 <210> SEO ID NO: 5 204 <211> LENGTH: 17

205 <212> TYPE: PRT

206 <213> ORGANISM: Artificial sequence

W--> 207 <220> FEATURE:

208 <223> OTHER INFORMATION: LSA-1 Consensus sequence of 17 amino acid repeats 209 where Xaa at position 1 is either Glu or Gly; Xaa at

position 4 is Ser or Arg; Xaa at position 6 is Asp or Ser; 210 Xaa at position 9 is Glu or Asp; Xaa at position 11 is Leu

-> 211/210) (W--> 212(2) or Arg; Xaa at position 13 is Lys or Asn and Xaa at position W--> 213 15 is Lys or Thr or Arg

215 <400> SEQUENCE: 5

W--> 216 Xaa Gln Gln Xaa Asp Xaa Glu Gln Xaa Arg 217 5

15

W--> 219 Xaa Ala Xaa Glu Xaa Leu Gln

E--> 220

VARIABLE LOCATION SUMMARY

PATENT APPLICATION: US/10/706,435A

DATE: 09/27/2004 TIME: 11:39:10

Input Set : A:\PTO.LM.txt

Output Set: N:\CRF4\09272004\J706435A.raw

Use of n's or Xaa's (NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing. Use of <220> to <223> is MANDATORY if n's or Xaa's are present. in <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

Seq#:5; Xaa Pos. 1,4,6,9,11,18,15

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/706,435A

DATE: 09/27/2004 TIME: 11:39:10

Input Set : A:\PTO.LM.txt

Output Set: N:\CRF4\09272004\J706435A.raw

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L:33 M:283 W: Missing Blank Line separator, <220> field identifier
L:35 M:283 W: Missing Blank Line separator, <400> field identifier
L:46 M:283 W: Missing Blank Line separator, <220> field identifier
L:48 M:283 W: Missing Blank Line separator, <400> field identifier
L:59 M:283 W: Missing Blank Line separator, <220> field identifier
L:61 M:283 W: Missing Blank Line separator, <400> field identifier
L:104 M:283 W: Missing Blank Line separator, <220> field identifier
L:106 M:283 W: Missing Blank Line separator, <400> field identifier
L:207 M:283 W: Missing Blank Line separator, <220> field identifier
L:212 M:259 W: Allowed number of lines exceeded, <223> Other Information:
L:213 M:259 W: Allowed number of lines exceeded, <223> Other Information:
L:216 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:5
L:216 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:5
L:216 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0
L:219 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:10
L:220 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:5
L:226 M:283 W: Missing Blank Line separator, <220> field identifier
L:228 M:283 W: Missing Blank Line separator, <400> field identifier
L:239 M:283 W: Missing Blank Line separator, <220> field identifier
L:241 M:283 W: Missing Blank Line separator, <400> field identifier
L:251 M:283 W: Missing Blank Line separator, <220> field identifier
L:253 M:283 W: Missing Blank Line separator, <400> field identifier
L:263 M:283 W: Missing Blank Line separator, <220> field identifier
L:265 M:283 W: Missing Blank Line separator, <400> field identifier
L:277 M:283 W: Missing Blank Line separator, <220> field identifier
L:279 M:283 W: Missing Blank Line separator, <400> field identifier
L:292 M:283 W: Missing Blank Line separator, <220> field identifier
L:294 M:283 W: Missing Blank Line separator, <400> field identifier
L:302 M:283 W: Missing Blank Line separator, <220> field identifier
L:304 M:283 W: Missing Blank Line separator, <400> field identifier
L:315 M:283 W: Missing Blank Line separator, <220> field identifier
L:317 M:283 W: Missing Blank Line separator, <400> field identifier
L:325 M:283 W: Missing Blank Line separator, <220> field identifier
L:327 M:283 W: Missing Blank Line separator, <400> field identifier
L:339 M:283 W: Missing Blank Line separator, <220> field identifier
L:341 M:283 W: Missing Blank Line separator, <400> field identifier
L:352 M:283 W: Missing Blank Line separator, <220> field identifier
L:354 M:283 W: Missing Blank Line separator, <400> field identifier
L:365 M:283 W: Missing Blank Line separator, <220> field identifier
L:367 M:283 W: Missing Blank Line separator, <400> field identifier
L:376 M:283 W: Missing Blank Line separator, <220> field identifier
L:378 M:283 W: Missing Blank Line separator, <400> field identifier
L:389 M:283 W: Missing Blank Line separator, <220> field identifier
L:391 M:283 W: Missing Blank Line separator, <400> field identifier
L:400 M:283 W: Missing Blank Line separator, <220> field identifier
L:402 M:283 W: Missing Blank Line separator, <400> field identifier
L:413 M:283 W: Missing Blank Line separator, <220> field identifier
L:415 M:283 W: Missing Blank Line separator, <400> field identifier
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VERIFICATION SUMMARY

PATENT APPLICATION: US/10/706,435A

DATE: 09/27/2004 TIME: 11:39:10

Input Set : A:\PTO.LM.txt

Output Set: N:\CRF4\09272004\J706435A.raw

```
L:425 M:283 W: Missing Blank Line separator, <220> field identifier L:427 M:283 W: Missing Blank Line separator, <400> field identifier L:438 M:283 W: Missing Blank Line separator, <220> field identifier L:440 M:283 W: Missing Blank Line separator, <400> field identifier L:448 M:283 W: Missing Blank Line separator, <220> field identifier L:4451 M:283 W: Missing Blank Line separator, <220> field identifier L:464 M:283 W: Missing Blank Line separator, <400> field identifier L:466 M:283 W: Missing Blank Line separator, <220> field identifier L:508 M:283 W: Missing Blank Line separator, <400> field identifier L:508 M:283 W: Missing Blank Line separator, <220> field identifier L:510 M:283 W: Missing Blank Line separator, <400> field identifier L:610 M:283 W: Missing Blank Line separator, <220> field identifier L:612 M:283 W: Missing Blank Line separator, <400> field identifier L:620 M:283 W: Missing Blank Line separator, <400> field identifier L:620 M:283 W: Missing Blank Line separator, <400> field identifier L:620 M:283 W: Missing Blank Line separator, <400> field identifier L:622 M:283 W: Missing Blank Line separator, <400> field identifier L:622 M:283 W: Missing Blank Line separator, <400> field identifier
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